

=> d his

(FILE 'HOME' ENTERED AT 11:08:35 ON 20 MAR 2005)

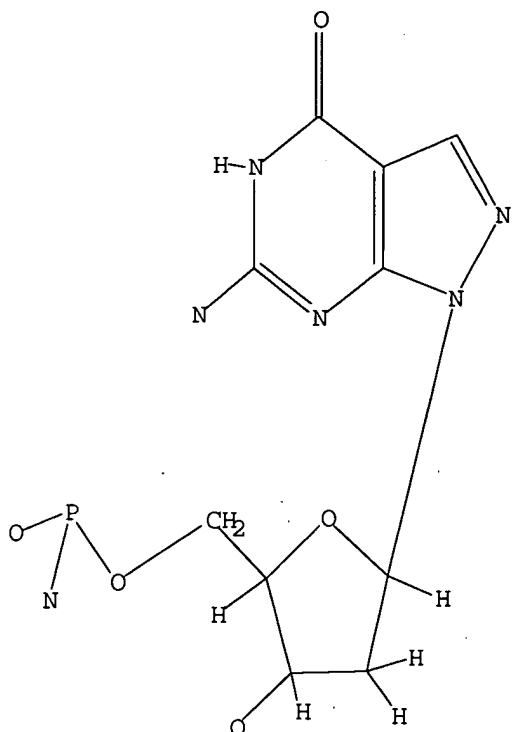
FILE 'REGISTRY' ENTERED AT 11:08:51 ON 20 MAR 2005

L1		STRUCTURE UPLOADED
L2	0	S L1 SSS SAM
L3	0	S L1 SSS FULL
L4		STRUCTURE UPLOADED
L5	6	S L4 SSS SAM
L6	77	S L4 SSS FULL
L7		STRUCTURE UPLOADED
L8	1	S L7 SSS SAM
L9	10	S L7 SSS FULL
L10		STRUCTURE UPLOADED
L11	1	S L10 SSS SAM
L12	9	S L10 SSS FULL
L13		STRUCTURE UPLOADED
L14	1	S L13 SSS SAM
L15	10	S L13 SSS FULL
L16		STRUCTURE UPLOADED
L17	0	S L16 SSS SAM
L18	0	S L16 SSS FULL

FILE 'CAPLUS, MEDLINE' ENTERED AT 11:43:26 ON 20 MAR 2005

L19	4	S L9 AND PHOSPHORAMIDITE?
L20	4	S L12 AND PHOSPHORAMIDITE?
L21	4	S L15 AND PHOSPHORAMIDITE?
L22	0	S L19 AND PHOTOLAB?
L23	0	S L20 AND PHOTOLAB?
L24	0	S L21 AND PHOTOLAB?

=> d 11
 L1 HAS NO ANSWERS
 L1 STR



Structure attributes must be viewed using STN Express query preparation.

=> s 11 sss sam
 SAMPLE SEARCH INITIATED 11:09:29 FILE 'REGISTRY'
 SAMPLE SCREEN SEARCH COMPLETED - 7 TO ITERATE

100.0% PROCESSED 7 ITERATIONS 0 ANSWERS
 SEARCH TIME: 00.00.01

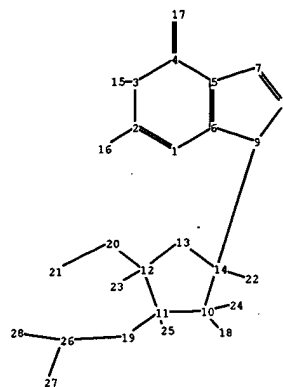
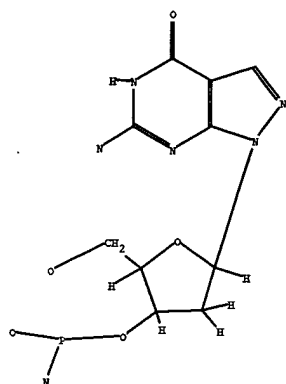
FULL FILE PROJECTIONS: ONLINE **COMPLETE**
 BATCH **COMPLETE**
 PROJECTED ITERATIONS: 7 TO 298
 PROJECTED ANSWERS: 0 TO 0

L2 0 SEA SSS SAM L1

=> s 11 sss full
 FULL SEARCH INITIATED 11:09:36 FILE 'REGISTRY'
 FULL SCREEN SEARCH COMPLETED - 75 TO ITERATE

100.0% PROCESSED 75 ITERATIONS 0 ANSWERS
 SEARCH TIME: 00.00.01

L3 0 SEA SSS FUL L1



chain nodes :

15 16 17 18 19 20 21 22 23 24 25 26 27 28

ring nodes :

1 2 3 4 5 6 7 8 9 10 11 12 13 14

chain bonds :

2-16 3-15 4-17 9-14 10-18 10-24 11-19 11-25 12-20 12-23 14-22 19-26 20-21
26-27 26-28

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 5-7 6-9 7-8 8-9 10-11 10-14 11-12 12-13 13-14

exact/norm bonds :

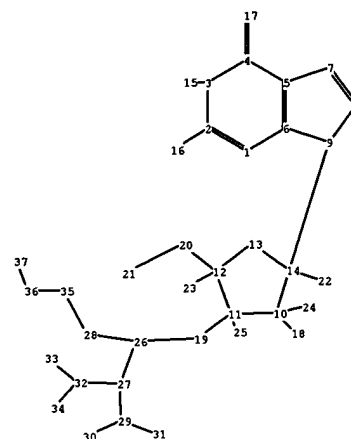
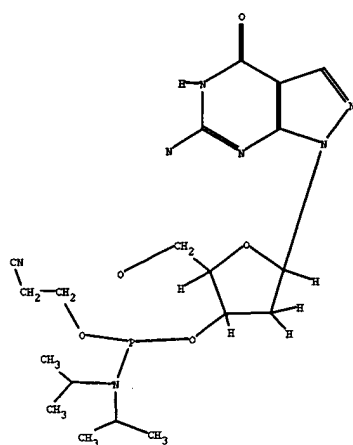
1-2 1-6 2-3 2-16 3-4 4-5 4-17 5-6 5-7 6-9 7-8 8-9 9-14 10-11 10-14 11-12
11-19 12-13 13-14 19-26 26-27 26-28

exact bonds :

3-15 10-18 10-24 11-25 12-20 12-23 14-22 20-21

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom 11:Atom
12:Atom 13:Atom 14:Atom 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS 20:CLASS
21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS



chain nodes :

15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36
37

ring nodes :

1 2 3 4 5 6 7 8 9 10 11 12 13 14

chain bonds :

2-16 3-15 4-17 9-14 10-18 10-24 11-19 11-25 12-20 12-23 14-22 19-26 20-21
26-27 26-28 27-29 27-32 28-35 29-30 29-31 32-33 32-34 35-36 36-37

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exact/norm bonds :

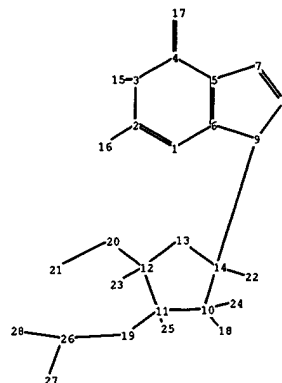
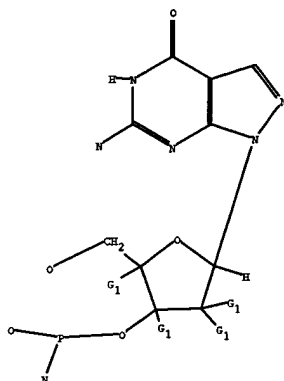
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35-36 36-37

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30:CLASS 31:CLASS 32:CLASS 33:CLASS 34:CLASS 35:CLASS 36:CLASS 37:CLASS



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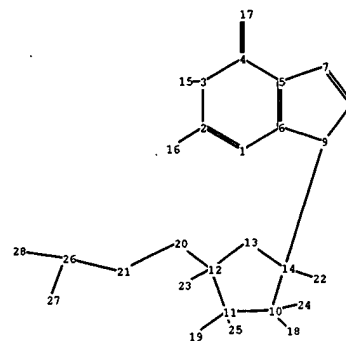
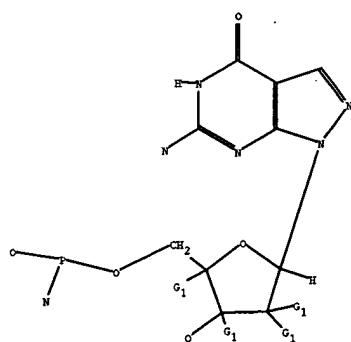
exact bonds :

3-15 12-20 14-22 20-21

G1:H,Cl,Br,F,I,O,Ak

Match level :

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